

PCT

WORLD INTELLECTUAL PROPERTY ORGANIZATION  
International Bureau

## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

<b>(51) International Patent Classification 5 :</b> <b>C07H 15/12, C12N 15/10</b> <b>C12P 19/34</b>		<b>A1</b> <b>(11) International Publication Number:</b> <b>WO 93/25563</b> <b>(43) International Publication Date:</b> 23 December 1993 (23.12.93)
<b>(21) International Application Number:</b> PCT/US92/05133 <b>(22) International Filing Date:</b> 17 June 1992 (17.06.92)		<b>(81) Designated States:</b> AU, CA, JP, US, European patent (AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LU, MC, NL, SE). <b>Published</b> <i>With international search report.</i>
<b>(71) Applicant (for all designated States except US):</b> CITY OF HOPE [US/US]; 1500 East Duarte Road, Duarte, CA 91010-0269 (US). <b>(72) Inventor; and</b> <b>(75) Inventor/Applicant (for US only) :</b> WALLACE, Robert, Bruce [CA/US]; 1020 Buena Vista, South Pasadena, CA 91030 (US). <b>(74) Agent:</b> IRONS, Edward, S.; 919 18th Street, N.W., Suite 800, Washington, DC 20006 (US).		
<b>(54) Title:</b> A METHOD OF DETECTING AND DISCRIMINATING BETWEEN NUCLEIC ACID SEQUENCES		
<b>(57) Abstract</b> <p>The present invention is directed to a method for detecting the presence or absence of any specific target nucleic acid sequence contained in a sample. The target sequence can be present in the sample in a relatively pure form or as a component of a member of a mixture of different nucleic acids. The method of the invention utilizes a novel primer design. The sequence of the novel primer is composed of two portions, the 3' portion is a primer specific for the desired nucleic acid sequence and the 5' portion is complementary to preselected nucleic acid sequence. Extension of the 3' portion of the primer with a labeled deoxynucleosides triphosphate yields a labeled extension product if, but only if, the template includes the target sequence. The labeled extension product is detected by hybridization of the 5' portion to the preselected sequence. The preselected sequence is preferably bound to a solid support as one member of a grid having a group of sequences.</p>		

***FOR THE PURPOSES OF INFORMATION ONLY***

Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

AT	Austria	GB	United Kingdom	MR	Mauritania
AU	Australia	GE	Georgia	MW	Malawi
BB	Barbados	GN	Guinea	NE	Niger
BE	Belgium	GR	Greece	NL	Netherlands
BF	Burkina Faso	HU	Hungary	NO	Norway
BG	Bulgaria	IE	Ireland	NZ	New Zealand
BJ	Benin	IT	Italy	PL	Poland
BR	Brazil	JP	Japan	PT	Portugal
BY	Belarus	KE	Kenya	RO	Romania
CA	Canada	KG	Kyrgyzstan	RU	Russian Federation
CF	Central African Republic	KP	Democratic People's Republic of Korea	SD	Sudan
CG	Congo	KR	Republic of Korea	SE	Sweden
CH	Switzerland	KZ	Kazakhstan	SI	Slovenia
CI	Côte d'Ivoire	LI	Liechtenstein	SK	Slovakia
CM	Cameroon	LK	Sri Lanka	SN	Senegal
CN	China	LU	Luxembourg	TD	Chad
CS	Czechoslovakia	LV	Latvia	TG	Togo
CZ	Czech Republic	MC	Monaco	TJ	Tajikistan
DE	Germany	MD	Republic of Moldova	TT	Trinidad and Tobago
DK	Denmark	MG	Madagascar	UA	Ukraine
ES	Spain	ML	Mali	US	United States of America
FI	Finland	MN	Mongolia	UZ	Uzbekistan
FR	France			VN	Viet Nam
GA	Gabon				

## REAGENTS COMPRISING CHIMERIC MOLECULES OF NUCLEIC ACIDS

## AND NUCLEIC ACID ANALOGS

\*\*\*\*\*

There are a series of compounds which have standard nucleic acid bases attached to nonstandard polymer backbones (these backbones can be nucleic acid

5 like; eg. phosphorothioate- or methylphosphonate-linked deoxyribose; or they can be very different; eg. polyamide). If the interbase spacing is correct, these compounds are capable of base pairing reactions analogous to normal nucleic acids. Interest in these  
10 compounds currently centres on their use in either therapeutics or diagnostic imaging.

The chemical and hybridization properties of some of these compounds are particularly interesting.

A number of novel and inventive uses of these

15 compounds as tools in Molecular Biology (as opposed to therapeutic and diagnostic pharmaceuticals as above) are presented herein. The synthesis and use of non-standard backboned nucleic acids has been described in WO 92/20702 and WO 92/20703. More recently WO 93/12129  
20 and WO 93/13121 described the synthesis and some uses of similar molecules.

In one aspect the present invention provides a chimeric nucleic acid/nucleic acid analogue molecule comprising a first portion, said first portion

25 comprising a nonstandard backboned oligonucleotide having at least one amide linkage and a second portion, said second portion comprising an acceptor end which is a chemical functionality capable of acting as acceptor for the formation of a phosphodiester bond.

30 The preferred backbone for the nonstandard backboned oligonucleotide is polyamide. The oligonucleotide preferably comprises peptide nucleic acid (PNA).

35 Preferably, the nonstandard backboned oligonucleotide will contain at least two monomers, and more preferably three or more monomers. These may be

- 2 -

monomers of PNA.

The chimeric molecules may be alternatively backboned nucleic acids that have a few normally backboned nucleotides at the 3' end. Such chimeric 5 molecules will combine the altered properties associated with the nonstandard backbone with an ability to prime DNA synthesis from a normal template-bound 3' end. Some of these alternatively backboned nucleic acids are capable of dsDNA strand invasion 10 below the target dsDNA melting temperature, somewhat like ssDNA/RecA complexes (Science, 254, p1497, (1991)).

Therefore new methods of sequencing are possible. This is particularly the case for dsDNA. A 15 suitable chimeric molecule is mixed with the dsDNA to be sequenced and then incubated with a DNA polymerase and nucleoside triphosphates and appropriate dideoxynucleoside triphosphate terminators. There should be no need to denature the dsDNA to be 20 sequenced. This is a major advantage especially for automated sequencing systems. In principle the 3' end of the sequencing primer may have any chemical functionality capable of acting as acceptor for the formation of a phosphodiester bond. This may be an 25 -OH group on the non-standard backboned molecule itself or a 3'-OH group if some normally backboned nucleotide(s) are required. Only one normally backboned nucleotide may be required but it is more likely that a length of 2 to 12 bases, more preferably 30 3 to 7 bases may be required. Work described in Biochemistry 29 1200-1207 (1990) shows that very short conventionally backboned primers may be utilised.

In addition to chimeric molecules as described herein, the invention also provides a method 35 of performing a primer extension reaction by the use

of

a) a target nucleic acid

b) a primer which is a chimeric nucleic acid/nucleic acid analogue molecule comprising a first

5 portion, said first portion comprising a nonstandard backboned oligonucleotide and a second portion, said second portion comprising an acceptor end which is a chemical functionality capable of acting as acceptor for the formation of a phosphodiester bond, said 10 chimeric molecule being capable of hybridising to part of the target

c) a supply of nucleotides

which method comprises mixing reagents a),

b) and c) in the presence of a chain extension enzyme

15 under conditions to allow the chimeric molecule to hybridise to the target and extension of the chimeric molecule at the acceptor end to occur, giving an extension product.

The invention also provides a method of

20 performing a chain termination reaction by the use of

a) a target nucleic acid

b) a chimeric nucleic acid/nucleic acid analogue molecule comprising a first portion, said first portion comprising a nonstandard backboned oligonucleotide and 25 a second portion, said second portion comprising an acceptor end which is a chemical functionality capable of acting as acceptor for the formation of a phosphodiester bond, said chimeric molecule being capable of hybridising to part of the target

c) a supply of nucleotides

d) a chain termination agent

which method comprises mixing reagents a), b)

c) and d) in the presence of a chain extension enzyme under conditions to allow the chimeric molecule to

35 hybridise to the target and extension of the chimeric molecule at the acceptor end to occur, so as to

produce terminated extension products, which terminated extension products are separated to allow part of the nucleotide sequence of the target nucleic acid to be determined.

5       A further aspect of the invention is a method of determining the nucleotide sequence of a target nucleic acid, which method comprises performing a chain termination method as described above, using a chain termination agent for each of the four different 10 nucleotides such that the nucleotide sequence of the target may be determined.

The preferred chimeric molecules for use as primers in the methods described herein are chimeric molecules discussed above according to the invention.

15      Thus, the primers preferably have at least one amide linkage in the backbone of the nonstandard backboned oligonucleotide.

20      Generally, a 6-mer is accepted as the smallest effective priming unit, although this can be reduced to 3 or 4 if very low temperatures are used. The chimeric molecule primers described herein are preferably 6 or more base units in length.

25      One or more of the reagents used in the various methods according to the invention may be labelled in ways which are known in the art.

There may be advantage in using an oligomer primer in which the majority (most or all) of the bases are pyrimidines. Such primers may prove to have enhanced strand invasion properties.

30      Whilst it will be appreciated that the use of chimeric molecules as herein described in Sanger chain-terminating sequencing is desirable, they may also be used in other related methods such as cycle sequencing.

35      The possibilities for using the chimeric molecules according to the invention and the

- 5 -

advantages these molecules provide in use also apply to other techniques including amplification techniques eg. PCR. It should therefore be possible to perform isothermal amplification of nucleic acids using the 5 molecules herein described. Quantification of products produced by PCR reactions may also be improved by these molecules. In existing methods the dsDNA product of the PCR reaction is denatured and mixed with an immobilised capture probe. Some 10 reannealing of the PCR product is inevitable and these molecules will not be captured by the immobilised probe leading to a loss of sensitivity. Use of PNA molecules, for example, in the capture step will eliminate the need to denature the PCR product leading 15 to greater capture and increased sensitivity (PNA oligomers are peptide nucleic acid molecules comprising nucleic acid bases attached to a peptide backbone through a suitable linker and are described in detail in WO 92/20702 and WO 92/20703). The binding 20 may also be stronger. The immobilisation of the capture probe may also be easier to achieve using conventional methods used for proteins and peptides.

Annealing of chimeric molecules at higher temperatures to provide increased specificity and 25 greater sensitivity in primer extensions, PCR and sequencing reactions is expected.

A normally backboned nucleotide or oligonucleotide could also be attached by its 3' end to the other end of the non standard backboned nucleic 30 acid so as to provide a normally backboned 5' end and facilitate kinase labelling etc.

Non standard backboned molecules could also be used in sequencing methods based on primer walking and subsequent developments by Studier (Science 11 35 December 1992 p 1787) . This original method makes use of sequence information near the terminus of a

previously sequenced fragment of DNA to generate a new primer that will allow the next contiguous piece of DNA to be sequenced. This process can be repeated a number of times. One disadvantage of this approach, 5 especially in genome sequencing projects, is the time required to determine and synthesise the next appropriate primer. This is slow and expensive.

The method developed by Studier uses a pre-synthesised pool of all possible different hexamers 10 (approximately 4,000) of known sequence. These are combined to form 12mers or 18 mers to make the required primer. The hexamers can be ligated together on their complementary template or held together using a single stranded binding protein or just allowed to 15 hybridise adjacent to each other.

If smaller primer units could be used then the number requiring synthesis would be much smaller. For example if 5mers could be used then approximately 1,000 would be required resulting in significant cost 20 savings. Unfortunately, small oligonucleotides do not hybridise strongly. The strong hybridisation properties of, for example, PNA oligomers or other non-standard backboned molecules overcomes this problem. A combination of random PNA molecules and 25 random standard oligonucleotides can be used to derive more efficient primer-walking strategies. The use of PNA-DNA chimeras which would hybridise more efficiently and also be primers for DNA polymerases is also possible.

30 The following list indicates some of the main uses for the chimeric molecules described herein. It will be clear to those skilled in the art that many other uses exist.

1. Primers for polymerases (using chimeric molecules 35 with normally backboned nucleic acids giving the 3' end: priming could also occur from strand invasion

- 7 -

complexes formed by reaction with dsDNA targets, a strand displacing polymerase would be used for the extension). This can also be extended to random primer labelling of dsDNA molecules using chimeric primers and strand invasion.

- 5 2. In situ hybridization with no need to denature the sample.
3. Primers/Probes that do not require dsDNA denaturation.
- 10 4. Improved sequencing by hybridization with short oligos.
5. Double strand sequencing without denaturation.
6. Isothermal PCR and related PCR alternatives.
7. Random primer labelling without denaturation.
- 15 8. Better probes for primer extension mapping procedures.
9. New labels and probes for multiplex sequencing.
10. Novel adapters/linkers for cDNA cloning.

20

#### TERMINOLOGY:

By nucleic acid(s), we mean either DNA or RNA of any chain length which can be either wholly or partially single or double stranded unless otherwise specified.

- 25 25 By "normally backboned", we refer to phosphodiester linked deoxyribose (for DNA) or phosphodiester linked ribose (for RNA) as the backbone to which the base residues (A, C, G and T for DNA and A, C, G and U for RNA) are linked.

- 30 30 By "nonstandard backboned", we refer to any polymers other than the normal phosphodiester linked deoxyribose (for DNA) or phosphodiester linked ribose (for RNA) as the backbone to which the base residues (A, C, G and T for DNA and A, C, G and U for RNA) are linked. Unless otherwise stated, the only requirement

for the nonstandard backboned polymer is that the interbase spacings are suitable for the formation of appropriate hydrogen bonds (Watson and Crick or triple helical or Hoogsteen type) with a normally backboned 5 nucleic acid target. Examples of nonstandard backbones are phosphorothioate linked deoxyribose, phosphorothioate linked ribose, methylphosphonate linked deoxyribose, methylyphosphonate linked ribose and polyamide. It will be immediately obvious to one 10 skilled in the art that there are many other possible backbone polymers allowing the correct interbase spacing and that this allows for a number of different chemical and physical properties specific to the backbone moiety to be exploited whilst preserving the 15 ability to bind to a complementary nucleic acid base sequence by hydrogen bonding.

It is also to be understood herein that chimeric molecules comprising nonstandard backboned nucleic acids with normally backboned nucleic acid 20 ends, for ligation, priming, labelling and other such applications known to those skilled in the art, are also possible. Such chimeric molecules are to be included in the Claims wherever the term "nonstandard backboned" is used unless otherwise specified. In 25 chimeric molecules the normally backboned sequence may be directly linked to the non-standard backboned sequence. It is also possible to include a small linker group between the two sequences.

By hybridization, we mean the sequence 30 specific binding between a probe (with A, C, G and T residues or A, C, G and U residues attached to a normally backboned or nonstandard backboned polymer as specified) and a target nucleic acid. For a wholly or partially double stranded target, the sequence 35 specific binding may also occur in a double stranded region by a process referred to herein as "strand

- 9 -

invasion". Strand invasion is where the sequence specific binding of probe occurs under conditions in which the target strands do not normally separate from each other (for example at temperatures below the 5 melting temperature of the target in a given solvent at a given ionic strength). Strand invasion does not normally occur with normally backboned probes. The application of this strand invading property of some nonstandard backboned nucleic acid probes to improve 10 existing and create novel Molecular Biology applications is a major inventive step.

Figures 1 to 5 show HPLC traces for the DNA and PNA/DNA molecules synthesised in Example 6:

15 Figures 1a and 1b - 15-mers of DNA and PNA/DNA, respectively;  
Figures 2a and 2b - 12-mers of DNA and PNA/DNA, respectively;  
Figures 3a and 3b - 9-mers of DNA and PNA/DNA, respectively;  
20 Figures 4a and 4b - 7-mers of DNA and PNA/DNA, respectively;  
Figures 5a and 5b - 5-mers of DNA and PNA/DNA, respectively.

25

#### EXAMPLES

##### Example 1

30 Synthesis of chimeric molecule with single normal backboned base  
35 0.5mg of PNA<sub>154</sub>-NH<sub>2</sub> (H-CAT CTA GTG A-LysNH<sub>2</sub>), synthesised according to the methods disclosed in WO 92/20702 and WO 92/20703, was mixed with 0.1mg 5'-amino

- 10 -

thymidine in 0.02ml 33%DMSO, 100 mM Tris-HCl pH 7.4. The reaction was started by the addition of 0.05mg subaric acid bis(N-hydroxysuccinimide) ester and incubated at room temperature for 24 hours.

5

Example 2

Synthesis of chimeric molecule with longer standard  
backboned sequence

10

0.5mg  $5'$  H<sub>2</sub>NTCG CAC TGC ATC  $3'$  (standard backbone) in 0.25ml 100mM Tris-HCl pH 7.4 was mixed with 0.5mg subaric acid bis(N-hydroxysuccinimide) ester in DMSO for 5 minutes. The reactants were purified by gel 15 filtration, freeze dried and resuspended in 0.1ml 100mM Tris-HCl pH 7.4. 0.5mg PNA<sub>154</sub> in 0.025ml Tris-HCl pH 7.4 was added and incubated for 24 hours at room temperature.

20

Example 3

PNA-DNA PRIMER EXTENSION ASSAY - GENERALISED PROTOCOL

Annealing conditions

25

The PNA-DNA primer synthesised as in examples 1 and 2 is diluted, 0, 1/10 1/100 and 1/1000 in 50mM tris, pH 7.5 containing 50mM NaCl and 7mM MgCl<sub>2</sub>.

30

The control DNA 11mer primer is diluted to 10pmole/ $\mu$ l in the same buffer.

The PNA-DNA primer and the control primer are mixed with equal volumes of 2pmole/ $\mu$ l template oligo to give 1pmole/ $\mu$ l template concentration.

35

Annealing is performed by boiling the mixtures for 3 minutes and then leaving the solutions

- 11 -

to cool to room temperature over a period of approximately 1 hour .

Primer Extension Assay

5

The extension reactions are carried out containing the PNA-DNA or the DNA primed template with 200 $\mu$ M dATP, dGTP, dTTP and 20 $\mu$ Ci of alpha  $^{32}$ P dCTP with exonuclease free Klenow polymerase.

10

The reactions are incubated at 37°C for 20 minutes then dCTP is added to a final concentration of 200 $\mu$ M. The reactions are incubated for a further 10 minutes then terminated by the addition of EDTA. Free nucleotide is removed by spin column centrifugation.

15

The results can be analysed by conventional denaturing polyacrylamide electrophoresis.

20

25

30

35

**Example 4**

The synthesis of the thymine PNA monomer is outlined in scheme 1.

All solvents were of HPLC grade; DCM and pyridine was distilled over  $\text{CaH}_2$ ; MeOH was distilled over Mg and  $\text{I}_2$ ; EtOH was distilled over  $\text{CaO}$  then Mg and  $\text{I}_2$ ; DMF was of peptide synthesis grade and was not further purified; anhydrous acetonitrile was purchased from Applied Biosystems Inc. (ABI). All other chemicals were supplied by Aldrich.

$^1\text{H}$  and  $^{13}\text{C}$  n.m.r. were recorded on a Brucker 250AC and Brucker 200WP spectrometer. Positive ion Fast Atom Bombardment (FAB) mass spectra were recorded on a Kratos MS50TC spectrometer using a thioglycerol matrix. Oligonucleotide synthesis was carried out on an ABI 394 DNA synthesiser. PNA/DNA synthesis (detritylation and cleavage from the solid support) was carried out on an ABI 380B DNA synthesiser.

Flash chromatography was carried out using silica gel 60 (Fluka). Thin layer chromatography (tlc) was carried out on aluminium sheets, silica 60 F254, 0.2mm layer (Merck) using the following solvent systems; A=nBuOH/AcOH/H<sub>2</sub>O (3:1:1 v:v), B=DCM/EtOAc (1:1 v/v), C=DCM/MeOH (9/1 v/v). Products were visualised using ninhydrin (1% w/v in EtOH), with heating for 5 min. ,short wave UV lamp(254nm) or via iodine oxidation.

**1-Thyminylacetic acid (I)**

Thymine (10.0g, 79.3 mmoles) was dissolved in water (50ml) containing KOH (17.1g, 0.30 moles) at 40°C. A solution of  $\text{BrCH}_2\text{CO}_2\text{H}$  (16.5g, 1.5 eq.) in water (25ml) was added dropwise over 30 min. and the reaction mixture heated at 40°C for 2h. The solution was cooled to room temperature, the pH adjusted to 5.5 (c. HCl) then stored at -4°C for 2h. The precipitate formed was removed by filtration and the filtrate adjusted to pH 2 (c. HCl). The precipitate formed was isolated by filtration and dried over  $\text{P}_2\text{O}_5$  (12.5g, 86%).  $R_f$ =0.1(A), FAB MS 185 ( $\text{M}+1$ )<sup>+</sup>,  $^1\text{H}$  n.m.r. (200MHz, DMSO); 1.74(s,3H,Me), 4.36(s,2H,CH<sub>2</sub>), 7.48(s,1H,Ar-H), 11.4(s,1H,NH).  $^{13}\text{C}$  n.m.r. (50MHz, DMSO); 12.1(Me), 48.7(CH<sub>2</sub>), 108.6(C5), 142.1(C6), 151.2(C2), 164.7(C4), 169.9(CO<sub>2</sub>H).

- 13 -

### 3-t-butoxycarbonylamino-1,2-propanediol (II)

3-amino-1,2-propanediol (10g, 0.11moles) was dissolved in water (250mL). The solution was cooled to 0°C and Boc anhydride (25g, 0.12 moles) added in one portion. The reaction mixture was brought to room temperature and the pH maintained at 10.5 using NaOH (2M) until the pH remained constant. The reaction mixture was stirred for a further 1h at room temperature then EtOAc (200mL) added and the pH adjusted to 2.5 using HCl (4M) at 0°C. The phases were separated and the aqueous phase extracted with EtOAc (10x150mL). The combined organic phases were dried ( $\text{MgSO}_4$ ) then evaporated *in vacuo* to yield V as an oil. The product solidified upon freezing (20.4g, 97%).  $R_f=0.75(\text{A})$ , 0.60(B), FAB MS 192 ( $\text{M}+1$ )<sup>+</sup>, <sup>1</sup>H n.m.r. (200MHz,  $\text{CDCl}_3$ ); 1.39(s,9H,Boc Me), 3.14(broad, 2H,  $\text{CH}_2$ ), 3.52(m,2H, $\text{CH}_2$ ), 3.68(m,1H,CH), 4.75(s,2H,2xOH), 5.52(m,1H,NH). <sup>13</sup>C n.m.r. (50MHz,  $\text{CDCl}_3$ ); 28.17(Me), 42.55(NCH<sub>2</sub>), 63.48(CH<sub>2</sub>), 71.08(CH), 79.76(quaternary C), 157.1(C=O).

### t-butoxycarbonylaminoacetaldehyde (III)

The diol V (5g, 26.1mmoles) was dissolved in water (50mL) and  $\text{NaIO}_4$  (6.8g, 1.2eq.) added. The reaction mixture was stirred at room temperature for 3h., filtered, then the aqueous solution extracted with DCM (5x100mL). The combined organic phases were dried ( $\text{MgSO}_4$ ) then evaporated to dryness *in vacuo* to yield the aldehyde as a colourless oil (3.9g, 94%). The product was used without further purification in the next step.  $R_f=0.7(\text{B})$ , 0.15(A); <sup>1</sup>H n.m.r. (200MHz,  $\text{CDCl}_3$ ); 1.38(s,9H,Boc), 3.98(s,2H, $\text{CH}_2$ ), 5.33(b,1H,NH), 9.56(s,1H,CHO). <sup>13</sup>C n.m.r (50MHz,  $\text{CDCl}_3$ ); 28.0 (Me), 51.1(CH<sub>2</sub>), 79.9(qC), 155.6(Boc C=O), 197.4(CHO).

- 14 -

### N-(t-butoxycarbonylaminoethyl)-glycine ethyl ester (IV)

The aldehyde III (5.13g, 32.2 mmoles) was dissolved in MeOH (100ml) and added to a mixture of ethyl glycinate (HCl salt, 11.2g, 2.5eq.) and Na BH<sub>3</sub>CN (2.02g, 1eq.). The reaction mixture was stirred at ambient temperature for 16h. then the solvent removed *in vacuo*. The residue was dissolved in water (100ml) and the pH adjusted to 8.0 using NaOH (2M). The aqueous phase was extracted with DCM (5x150ml), the combined organic fractions dried (MgSO<sub>4</sub>), then evaporated *in vacuo* to an oil which was purified by Kugelrohr distillation (6.92g, 87%) R<sub>f</sub>=0.55(B), 0.15(C),  
<sup>1</sup>H n.m.r. (250MHz, CDCl<sub>3</sub>); 1.09(t,3H,CH<sub>3</sub>),  
1.25(s,9H,BocMe), 2.53(t,2H,CH<sub>2</sub>), 2.90(b,1H,NH), 3.46(t,2H,CH<sub>2</sub>NH),  
3.54(s,2H,CH<sub>2</sub>), 4.00(q,2H,CH<sub>2</sub>CH<sub>3</sub>), 5.46(b,1H,carbamateNH).  
<sup>13</sup>C n.m.r.(50MHz, CDCl<sub>3</sub>); 13.69(CH<sub>3</sub>CH<sub>2</sub>), 27.91(BocMe), 39.6, 42.6,  
48.27, 60.36(all CH<sub>2</sub>), 78.58(quat. C), 156.1(BocC=O), 172.4(C=O).

### N-(t-butoxycarbonylaminoethyl)-N-(thyminylacetyl)-glycine-ethyl ester (V)

Bocaminoethylglycine ethyl ester (IV, 2.75g, 11mmoles) was dissolved in DMF (12mL) and thyminylacetic acid (I, 2.05g, 1eq.) was added. On dissolution of the acid DCM (10mL) was added, the reaction mixture cooled to 0°C and DCC (2.49g, 1.2eq.) added. The reaction was stirred at 0°C for 1h. then for a further 2h. at room temperature. The precipitated DCU was removed by filtration, washed with DCM (2x30mL) and a further volume of DCM (150mL) added. The combined organic phases were washed with NaHCO<sub>3</sub> (0.5M, 3x100mL), citric acid (10%w/v, 2x100mL), and NaCl solution (sat., 1x100mL), dried (MgSO<sub>4</sub>) then evaporated to dryness *in vacuo* to yield a brown oil. The crude product was purified by silica gel chromatography eluting with DCM/MeOH(1-3%) to yield the product as a white foam (1.86g, 41%), R<sub>f</sub>=0.55(A),0.15(C) FAB MS 413 (M+1)<sup>+</sup>, <sup>1</sup>H n.m.r. (250MHz, CDCl<sub>3</sub>); (some of the signals are split into major and minor peaks due to restricted rotation around the secondary amide bond.) 1.18(t,3H,MeCH<sub>2</sub>),  
1.35(s,9H,BocMe), 1.81(s,3H,TMe), 3.25(m,2H,CH<sub>2</sub>), 3.50(m,2H,CH<sub>2</sub>),

- 15 -

4.00(s,2H,NCH<sub>2</sub>CO<sub>2</sub>Et), 4.39(s,mi,CH<sub>2</sub>CON), 4.53(s,mj,CH<sub>2</sub>CON), 5.76(b,1H,BocHN), 6.99(s,mj,Ar-H), 7.0(s,mi,Ar-H), 10.0(s,mj,NH), 10.05(s,mi,NH). <sup>13</sup>C n.m.r. (50MHz,CDCl<sub>3</sub>); 12.1(CH<sub>3</sub>CH<sub>2</sub>), 13.8(T-Me), 28.14(BocMe), 38.4, 47.6, 48.4, 48.7, 61.4(all CH<sub>2</sub>), 79.6(quat.C), 140.84(C6), 151.0, 155.9, 164.4, 167.2, 169.4(all C=O).

**N-(Monomethoxytritylaminooethyl)-N-(thyminylacetyl)-glycine-ethyl ester (VI)**

1-(Bocaminoethylglycine)-thymine ethyl ester (VII, 2.0g, 4.85mmoles) was dissolved in TFA (10mL). The reaction mixture was evaporated to dryness *in vacuo* then the residue coevaporated with toluene (5x10ml) then pyridine (2x10mL). Monomethoxytrityl chloride (1.65g, 1.1eq.), 4-pyrrolidinylpyridine (25mg) and pyridine (10mL) were added and the solvent was removed *in vacuo* once more. Pyridine (10mL) was added and the reaction mixture was stirred for 16h. Water (30mL) was added and the reaction mixture extracted with DCM (3x70mL). The combined organic phases were dried (MgSO<sub>4</sub>), then evaporated to dryness *in vacuo* to yield an oil. The product was isolated by column chromatography using DCM/MeOH(0-3%) as the eluant (2.55g, 90%). The SiO<sub>2</sub> was pre-equilibrated with DCM containing NEt<sub>3</sub> (1%v/v) to prevent detritylation. R<sub>f</sub>=0.2(A), 0.1(A); FAB MS 585 (M+1)<sup>+</sup>, <sup>1</sup>H n.m.r. (250MHz, CDCl<sub>3</sub>); (some of the signals were split into major and minor peaks due to restricted rotation about the secondary amide bond) 1.18(t,3H,MeCH<sub>2</sub>), 1.90(s,3H,T-Me), 2.38(m,2H,CH<sub>2</sub>NCO), 3.52(m,2H,TrNH-CH<sub>2</sub>), 3.76(s,3H,OMe), 3.90(s,mj,CH<sub>2</sub>CON), 4.02(s,mi,CH<sub>2</sub>CON), 4.10(q,2H,CH<sub>2</sub>Me), 4.25(s,mj,CH<sub>2</sub>CO<sub>2</sub>Et), 4.31(s,mi,CH<sub>2</sub>CO<sub>2</sub>Et), 6.82(s,mi,H6), 6.85(s,mj,H6), 7.2-7.4(m,Ar-H). <sup>13</sup>C n.m.r.(50MHz, CDCl<sub>3</sub>); 12.2(CH<sub>3</sub>CH<sub>2</sub>), 13.9(T-Me), 41.7, 48.0, 48.5, 50.0, 61.3(all CH<sub>2</sub>), 55.0(OMe), 110(CPh<sub>3</sub>), 123-128(m,Ar-H), 137.4(quatC), 140.8(C6), 145.6(quatC), 149.5(quatC), 151.0(COMe), 157.8, 164.1, 167.3, 168.8(allC=O).

**N-(Monomethoxytritylaminoethyl)-N-(thyminylacetyl)-glycine-(VII)**

The ester XI (1.34g, 2.3mmoles) was dissolved in MeOH (60mL) and NaOH (2M, 40mL) added. The reaction mixture was stirred at room temperature for 2h. then DOWEX (pyridinium form) was added until a pH of 7 was obtained. The suspension was filtered and the resin washed with MeOH (3x50mL). The filtrate was evaporated to dryness *in vacuo* to yield XI (1.4g, 96%).  $R_f$ =0.05(B),0(C);  $^1H$  n.m.r. (200MHz,CDCl<sub>3</sub>); (some of the signals were split into major and minor peaks due to restricted rotation about the secondary amide bond) 1.75(s,3H,T-Me), 2.21(m,2H,CH<sub>2</sub>), 3.49(m,2H,CH<sub>2</sub>), 3.76(s,3H,OMe), 3.85(s,mj,CH<sub>2</sub>CON), 4.05(s,mi,CH<sub>2</sub>CON), 4.52(s,mj,CH<sub>2</sub>CO<sub>2</sub>H), 4.85(s,mi,CH<sub>2</sub>CO<sub>2</sub>H), 6.85(s,mj,H6), 6.90(s,mi,H6), 7.2-7.5(m,Ar-H).  $^{13}C$  n.m.r.(50MHz,CDCl<sub>3</sub>); 12.08(T-Me), 33.48, 42.15, 48.23, 48.38 (all CH<sub>2</sub>), 55.07(OMe), 113.2(CPh<sub>3</sub>), 126-131(m,Ar-H), 137.4, 146.0, 149.8 (all quat C), 150.8(COMe), 158.9, 164.4, 168.3, 173.3(C=O).

Example 5

The nucleoside 5'-monomethoxytritylamino-5'-deoxy thymidine phosphoramidite was synthesised as shown in scheme 2 and was used as the initial linker in the PNA DNA chimeric molecule.

**5'-azido-5'deoxy thymidine (VIII)**

Thymidine (14.0g, 58.0mmoles) was stirred in DMF (175ml) and PPh<sub>3</sub> (18.2g,1.2eq.), NaN<sub>3</sub> (11.3g, 3eq.), and CBr<sub>4</sub> (23.0g, 1.2eq.) added at 0°C. The reaction mixture was stirred at 0°C for 1h. then ambient temperature for 16h. The reaction mixture was poured onto NaHCO<sub>3</sub> (0.5M, 350ml) and extracted with DCM (6x150ml). The combined organic phases were dried (MgSO<sub>4</sub>) then evaporated to an oil which was purified by SiO<sub>2</sub> chromatography (0-4% MeOH/DCM , 7.73g, 50%).  $R_f$ =0.1(C), 0.8(A), FABMS (M+1) 268,  $^1H$  n.m.r. (200MHz, D6-DMSO) 1.78(s,3H,T-CH<sub>3</sub>), 2.10(m,1H,2'H), 2.26(b,1H,2'H), 3.56(d,2H, 5'H), 3.85(m,1H,4'H), 4.20(m,1H, 3'H), 5.42(d,1H,3'OH), 6.21(t,1H,1'H)

- 17 -

7.5(s,1H,6H), 11.34(s,1H,NH).  $^{13}\text{C}$  n.m.r.(50MHz, D6-DMSO); 12.2(CH<sub>3</sub>), 33.9(CH<sub>2</sub>), 51.8(CH<sub>2</sub>), 70.9,72.1(CH), 136.2(C6), 150.6, 163.8(C=O).

### 5'-N-(4-Methoxytrityl)amino-5'-deoxythymidine (X)

To the azido compound (7.73g, 29mmoles) in pyridine (100ml) was added PPh<sub>3</sub> (15.2g, 1.2 eq.). The reaction mixture was stirred at room temperature for 16h. then water (20ml) added. After stirring for 1h. the reaction mixture was poured onto water (300ml) and stirred for a further 16h. The suspension was filtered and the aqueous phase washed with EtOAc (3x150ml). The 5'-amino-5'-deoxy-thymidine was isolated by freeze drying (4.1g, 58%).

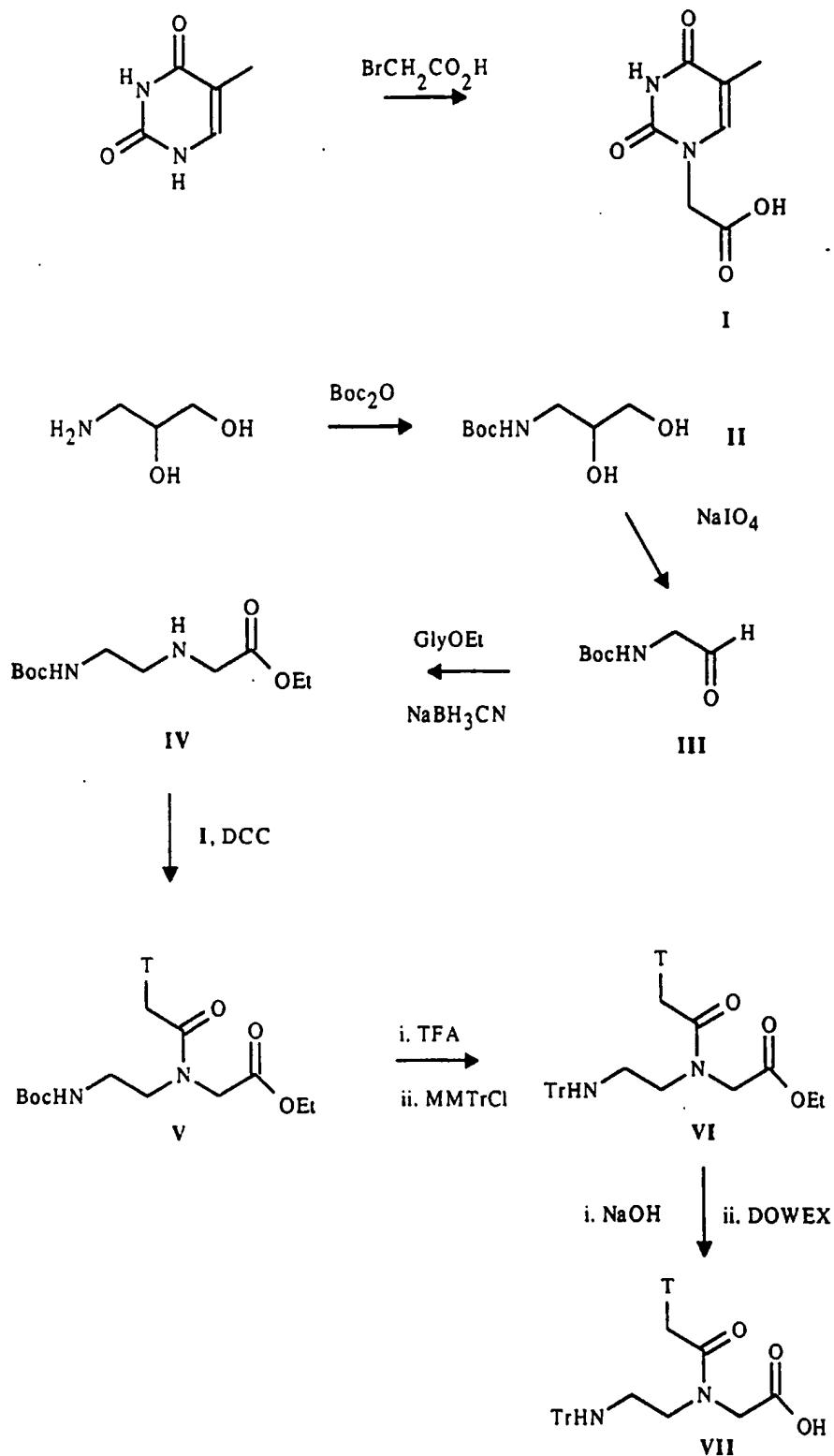
A fraction of the 5'-amino-5'-deoxy-thymidine (1.34g, 5.55 mmole) was coevaporated with pyridine (2x20ml) then dissolved in pyridine (40ml). 4-pyrollidinopyridine (30mg) and p-anisoyldiphenylmethyl chloride (2.05g, 1.2eq.) were added, the reaction mixture stirred at room temperature for 16h. then poured onto NaHCO<sub>3</sub> solution (0.5M,30ml). The aqueous phase was extracted with DCM (3x50ml), the combined organic phases dried (MgSO<sub>4</sub>) then evaporated to dryness *in vacuo*. The resultant oil was purified by column chromatography (SiO<sub>2</sub> preequilibrated with 1% NEt<sub>3</sub> in DCM eluting with 0-3% MeOH/DCM). (2.02g,71%) R<sub>f</sub>=0.5(C), 0.2(B). FABMS (M+1)=514  $^1\text{H}$  n.m.r. (200MHz, D6-DMSO); 1.69(d,3H,CH<sub>3</sub>), 2.01-2.34(m,4H,2'H,5'H), 2.64(t,1H,5'NH), 3.72(s,3H,OMe), 3.83(m,1H,4'H), 4.19(m,1H,3'H), 5.22(d,1H,3'-OH), 6.14(t,1H,1'H), 6.84 (d,2H,OMe), 6.87-7.43(m,13H,aromatic), 11.29(s,1H,NH).  $^{13}\text{C}$  n.m.r.(50MHz, D6-DMSO); 12.2(CH<sub>3</sub>), 24.9(CH<sub>2</sub>), 40.0(CH<sub>2</sub>), 54.8(OMe), 71.6(CH), 84.0(CH), 85.9(CH), 112.8-129.4(aromatic CH), 134.9, 137.4, 145.6, 147.8, 157.5(all qC), 150.4(C=O), 164.1(C=O).

- 18 -

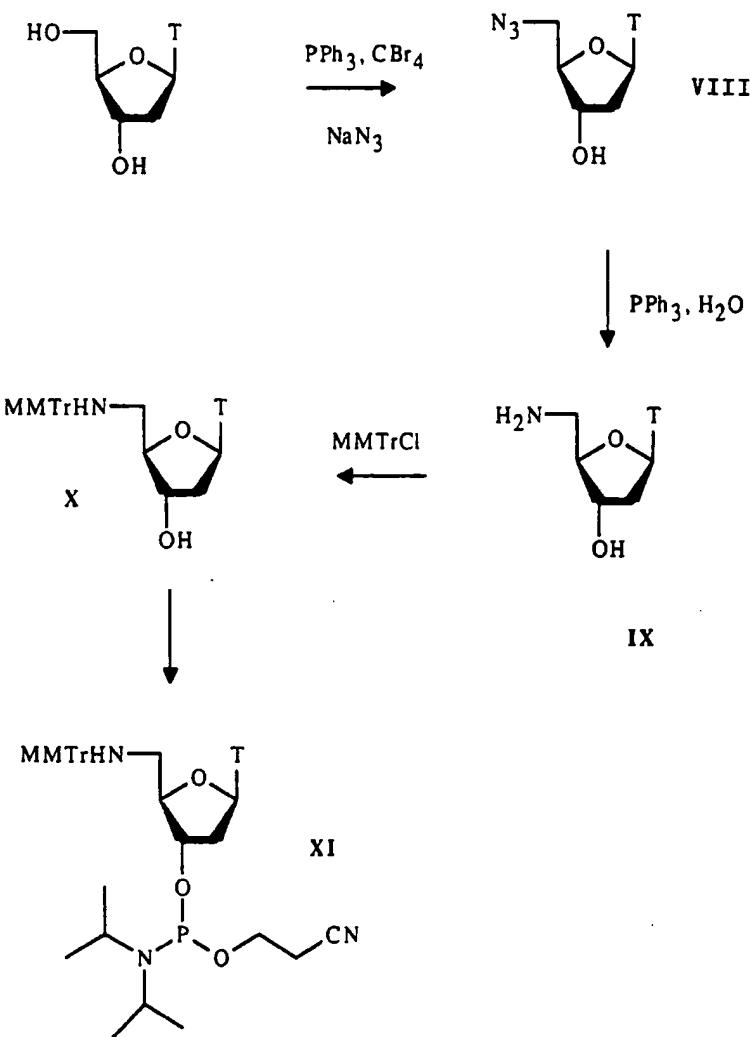
**5'-N-(4-Methoxytrityl)amino-5'-deoxythymidine-3'-O-(2-cyanoethyl,  
N,N-diisopropylamino)phosphite (XI)**

5'-N-(4-Methoxytrityl)amino-5'-deoxythymidine (0.44g, 0.85mmoles) and diisopropylammonium tetrazolide (30mg, 0.25eq.) were coevaporated with pyridine (2x5ml) then MeCN (5ml). The residue was dissolved in MeCN (5ml) then 2-cyanoethyl-N,N,N',N'-tetraisopropylphosphorodiamide (0.35ml, 1.4eq.) added and the reaction mixture stirred at room temperature for 16h. DCM (50ml) was added and the solution washed with NaHCO<sub>3</sub> (0.5M, 20ml), dried (MgSO<sub>4</sub>) then evaporated to an oil which was purified by SiO<sub>2</sub> chromatography (100% EtOAc, 0.30g, 49%).  
R<sub>f</sub>=0.77 (100% EtOAc).

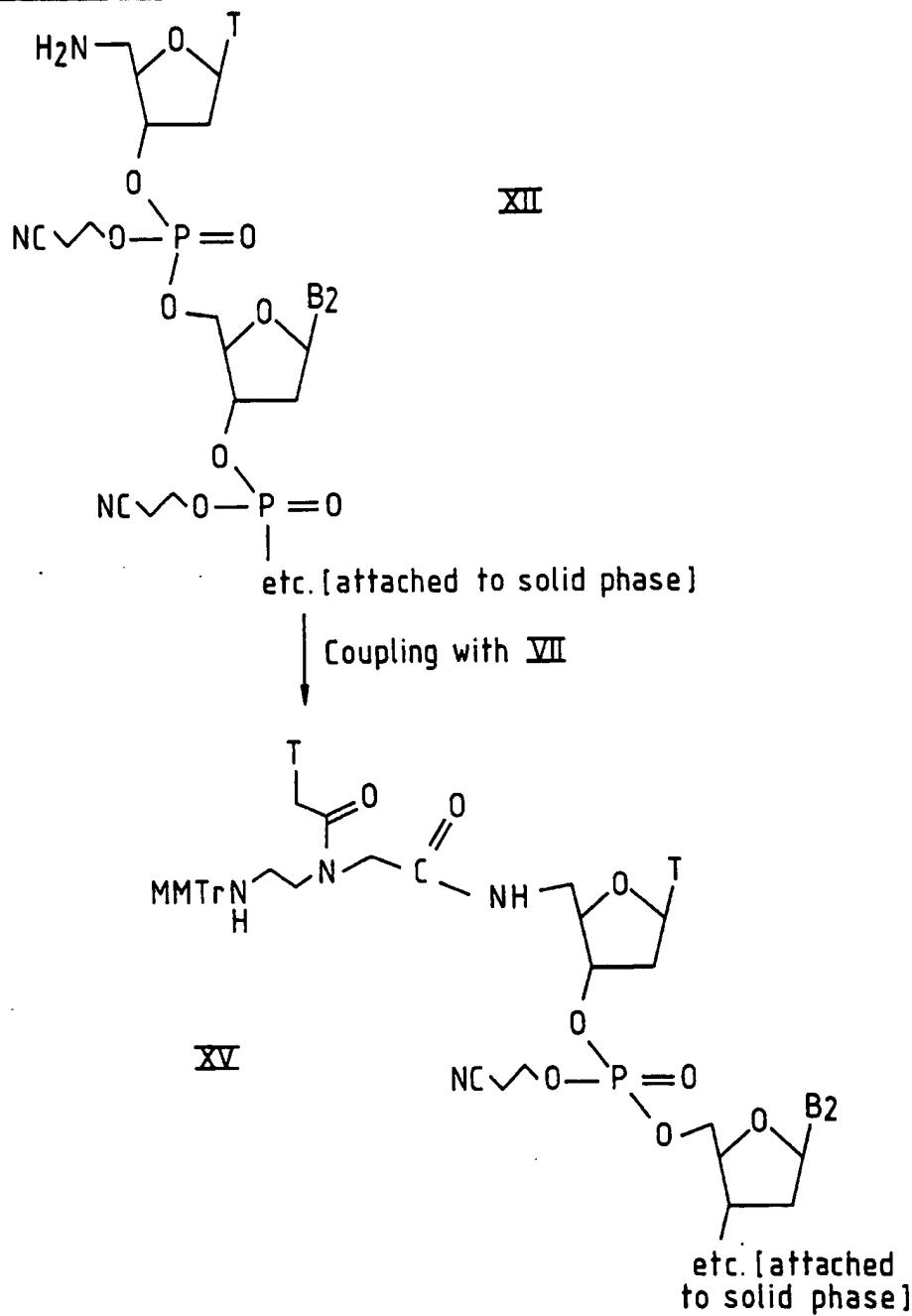
- 19 -

Reaction Scheme 1

- 20 -

Reaction Scheme 2

- 21 -

Reaction Scheme 3

Example 6Solid Phase Synthesis of PNA/DNA Chimeric Compounds

5 Chimeric molecules were made by initial synthesis of the DNA moiety using standard phosphoramidite chemistry and standard automated synthesis on an ABI 394 synthesiser. The final DNA monomer addition was a modified nucleoside  
10 phosphoramidite; a 5'-amino-5'-2'-dideoxy nucleoside derivative, protected at the 5'-end with a monomethoxytrityl group (XI in Reaction Scheme 2). This was deprotected at the 5'-end with trichloroacetic acid in the standard way and the solid phase bound  
15 oligonucleotide was then ready for the addition of the PNA monomers. The PNA section of the molecule was then added in a stepwise fashion using a manual solid phase methodology.

20 Standard procedure for manual solid phase coupling

Reaction Scheme 3 shows coupling of support bound DNA with PNA.

The trityl protecting group of the support  
25 bound DNA was cleaved using the appropriate cycle on an ABI 380B DNA synthesiser giving solid phase DNA with a 5'-amino group ready for a PNA coupling reaction XII. The PNA monomer VII (which is protected with a monomethoxy trityl group at the amino terminus) 280 mg  
30 was dissolved in DMF/Pyridine (1:1 v/v 4.4ml) to yield a 0.1M solution. An aliquot (0.1ml) was added to the peptide coupling reagent HBTU (18mg, 5eq.), and DECHA (20 $\mu$ l, 10 eq.) then manually passed repeatedly through the column containing the supported DNA for 1h. The  
35 coupling procedure was then repeated, to improve yield. The PNA monomethoxytrityl group was then removed on the

- 23 -

solid phase using trichloroacetic acid and the next PNA coupling reaction was carried out as previously.

When the desired number of monomers had been added, the molecule was cleaved from the solid support using a standard cycle on the DNA synthesiser and the protecting groups removed by heating in  $\text{NH}_3$  at 55°C for 5h. The product was purified by reverse phase HPLC. HPLC purification was carried out on a Gilson 303 system using a reverse phase octadecyl stationery phase. The PNA/DNA molecules were purified "trityl off" and the buffer system employed was; A= $\text{NH}_4\text{OAc}$  (0.1M), B= MeCN (30% v/v)/ $\text{NH}_4\text{OAc}$  (0.1M). The flow rate was 3ml/min, detector 280nm.

The PNA/DNA chimeric molecules in Table 1 below were synthesised according to the above method. PNA moieties appear in lower case and DNA bases in upper case. Equivalent DNA sequences with standard T nucleotides were also synthesised. Figures 1 to 5 show compared HPLC data for the PNA/DNA and DNA sequences. Clear peaks were obtained and in all cases mobility of PNA/DNA molecules was retarded compared to DNA molecules of equivalent length and sequence.

Table 1

25	SIZE	SEQUENCE	FIGURE NO.
	15-mer	tttTAGAGTGTTGTT	1b
	12-mer	tttTAGAGTGTT	2b
	9-mer	tttTAGAGT	3b
30	7-mer	tttTAGA	4b
	5-mer	tttTA	5b

Example 7PNA-DNA Primer Extension Assay5 Annealing Conditions

The 18-mer DNA template comprised the following sequence:

10 5'-CTGAACAAACACTCTAAAA-3'

The PNA-DNA primers were synthesised as described in example 6 and shown in Table 1 above.

15 The control primers comprised DNA only, having base sequences identical to the PNA-DNA primers described.

Annealing reactions were carried out in a total volume of 10 $\mu$ l in the presence of half strength Klenow buffer  
20 (1X Klenow buffer contains: 50mM Tris-HCl (pH7.5), 5mM MgCl<sub>2</sub>, 5mM  $\beta$ -Mercaptoethanol). The concentrations of primer and template were 5 and 1 pmol respectively.

Annealing was performed by boiling the reaction  
25 mixtures for 3 minutes and then leaving the samples at room temperature for 30 minutes.

Primer Extension Assay

30 The extension reactions contained the contents of the annealing reactions described above (10 $\mu$ l each) together with 20 $\mu$ M dGTP, 20 $\mu$ M dATP, 20 $\mu$ M dTTP, 2.5 $\mu$ Ci  $\alpha$ -<sup>32</sup>PdCTP (3000Ci/mmol, Amersham), 1.5X Klenow buffer, Klenow or exonuclease free Klenow (United States  
35 Biochemical) at 1 unit per reaction in a total volume of

- 25 -

20 $\mu$ l. (Klenow is a fragment of the enzyme DNA polymerase 1).

5 5mM d ATP  $\alpha$ -S was included in reactions containing Klenow enzyme in order to prevent the 3'-5' exonuclease activity of the said enzyme.

Reactions were incubated at 37°C for up to 30 minutes. Reaction products were analysed by polycrylamide gel electrophoresis or TLC as described below.

10

Thin Layer Chromatography (TLC) Analysis System

15 2 $\mu$ l of each reaction mix was added to 3 $\mu$ l of stop solution (95% formamide, 20mM EDTA, 0.05% bromophenol blue and xylene cyanol FF). 1.0 $\mu$ l of each stopped reaction was then spotted onto a Schleicher and Schuell PEI-cellulose TLC plate 20mm up from the base of the plate. The plates were developed in a tank containing 1M KH<sub>2</sub>PO<sub>4</sub>. The plates were allowed to run until the 20 solvent front had reached 10mm from the top of the plate. The plates dried and analysed on a Raytest TLC scanner. Labelled reaction products stayed on or close to the origin whereas the [ $\alpha$ <sup>32</sup>P] dNTP or excised [<sup>32</sup>P] dNMP moved up the plate according to the base used and 25 the number of phosphates.

Denaturing Polyacrylamide gel electrophoresis (PAGE)

30 Page I<sup>TM</sup> Sequencing Gel Mix (19:1) from Boehringer Mannheim Corporation was diluted according as described by the manufacturer to give an 18% polyacrylamide gel mix containing 7M urea. 70ml of this mix was taken and polymerised by the addition of 22 $\mu$ l TEMED and 650 $\mu$ l 10% ammonium persulphate. This was used to pour a standard 35 0.4mm sequencing gel. The gel was run in 1x TBE buffer.

- 26 -

Gels were pre-run for 30 minutes at 1.5kV. Standard sharks tooth combs were used and 2.5 $\mu$ l of each sample containing 5000cpm/ $\mu$ l in 95% formamide, 20mM EDTA, 0.05% bromophenol blue and xylene cyanol FF was loaded 5 per well. Gels were run for 2 hours at 2.8kV. Gels were fixed in 10% acetic acid: 10% methanol and dried onto Whatman 3MM paper. The dried gel was then exposed to  $\beta$ max film (Amersham).

10

Results

Control experiments were set up using the DNA primers and 18mer template. The DNA oligo primer sizes were 15 4mer, 5mer, 7mer, 9mer, 12mer, & 15mer respectively. Only the 9, 12, and 15mer gave efficient priming of the template as followed by the incorporation of [ $\alpha$ <sup>32</sup>P]dNTP on TLC plates. The 7mer did prime the template allowing extension but to a lesser degree than the larger DNA 20 oligo's. This was true when both Klenow and exonuclease free Klenow enzymes were used respectively.

Further experiments used the PNA-DNA chimeric primers to prime the template, again priming of the template 25 was followed by the incorporation of [ $\alpha$ <sup>32</sup>P]dNTP when analysed by TLC. Only the PNA-DNA 5mer failed to show any incorporation of the radiolabel. All the other four chimeras showed efficient incorporation of the radiolabel in the presence of either Klenow and dATP $\alpha$ S 30 or exonuclease free Klenow.

The mobility of the labelled reaction products from the PNA-DNA chimera was significantly retarded and ran at a higher molecular weight compared to the corresponding 35 DNA primer reaction products in polyacrylamide gel electrophoresis.

- 27 -

Competition extension reactions involving the equivalent chimera and DNA primers in the presence of the template resulted in the chimera being preferentially radiolabelled and no radiolabelled DNA 5 product was produced.

Abbreviations

	DCM	dichloromethane
10	DCC	dicyclohexylcarbodiimide
	DCU	dicyclohexylurea
	DECHA	diethylcyclohexylamine
	DMF	dimethylformamide
	EtOAc	ethyl acetate
15	HBTU	hydroxybenzatriazolyltetramethyluroniumhexa-fluorophosphate
	MMTr	p-Anisoyldiphenylmethane
	HPLC	High Pressure Liquid Chromatography

20

25

30

35

CLAIMS

1. A chimeric nucleic acid/nucleic acid analogue  
5 molecule comprising a first portion, said first portion comprising a nonstandard backboned oligonucleotide having at least one amide linkage and a second portion, said second portion comprising an acceptor end which is a chemical functionality capable of acting as acceptor  
10 for the formation of a phosphodiester bond.
2. A chimeric molecule as claimed in claim 1, wherein the second portion comprises at least one normally backboned nucleotide and has a 3' end.
3. A chimeric molecule as claimed in claim 1 or  
15 claim 2, wherein the molecule is labelled.
4. A chimeric molecule as claimed in any one of claims 1 to 3, wherein the nonstandard backboned oligonucleotide has a polyamide backbone.
5. A chimeric molecule as claimed in claim 4,  
20 wherein the nonstandard backboned oligonucleotide is PNA.
6. A method of performing a primer extension, reaction by the use of
  - a) a target nucleic acid
  - 25 b) a primer which is a chimeric nucleic acid/nucleic acid analogue molecule comprising a first portion, said first portion comprising a nonstandard backboned oligonucleotide and a second portion, said second portion comprising an acceptor end which is a chemical functionality capable of acting as acceptor  
30 for the formation of a phosphodiester bond, said chimeric molecule being capable of hybridising to part of the target
  - c) a supply of nucleotides  
35 which method comprises mixing reagents a), b)

and c) in the presence of a chain extension enzyme under conditions to allow the chimeric molecule to hybridise to the target and extension of the chimeric molecule at the acceptor end to occur, giving an  
5 extension product.

7. A method as claimed in claim 6, wherein at least one of reagents a), b) and c) is labelled.

8. A method of amplifying a target nucleic acid by an amplification reaction, wherein at least one  
10 primer is used which is a chimeric nucleic acid/nucleic acid analogue molecule comprising a first portion, said first portion comprising a nonstandard backboned oligonucleotide and a second portion, said second portion comprising an acceptor end which is a  
15 chemical functionality capable of acting as acceptor for the formation of a phosphodiester bond.

9. A method as claimed in claim 8, wherein the amplification reaction is the polymerase chain reaction.

20 10. A method of performing a chain termination reaction by the use of  
a) a target nucleic acid  
b) a primer which is a chimeric nucleic acid/nucleic acid analogue molecule comprising a first portion, said first portion comprising a nonstandard backboned oligonucleotide and a second portion, said second portion comprising an acceptor end which is a chemical functionality capable of acting as acceptor for the formation of a phosphodiester bond, said  
25 chimeric molecule being capable of hybridising to part of the target  
30 c) a supply of nucleotides  
d) a chain termination agent

which method comprises mixing reagents a), b)

35 c) and d) in the presence of a chain extension enzyme under conditions to allow the chimeric molecule to

- 30 -

hybridise to the target and extension of the chimeric molecule at the acceptor end to occur, so as to produce terminated extension products, which terminated extension products are separated to allow

5 part of the nucleotide sequence of the target nucleic acid to be determined.

11. A method as claimed in claim 10, wherein at least one of the reagents b), c) and d) is labelled.

12. A method of determining the nucleotide 10 sequence of a target nucleic acid, which method comprises performing the method of claim 10 or claim 11 using a chain termination agent for each of the four different nucleotides such that the nucleotide sequence of the target may be determined.

15 13. A method as claimed in any one of claims 6 to 12, wherein the target nucleic acid is present in a double-stranded nucleic acid.

20

25

30

35

1/5

Fig.1a.

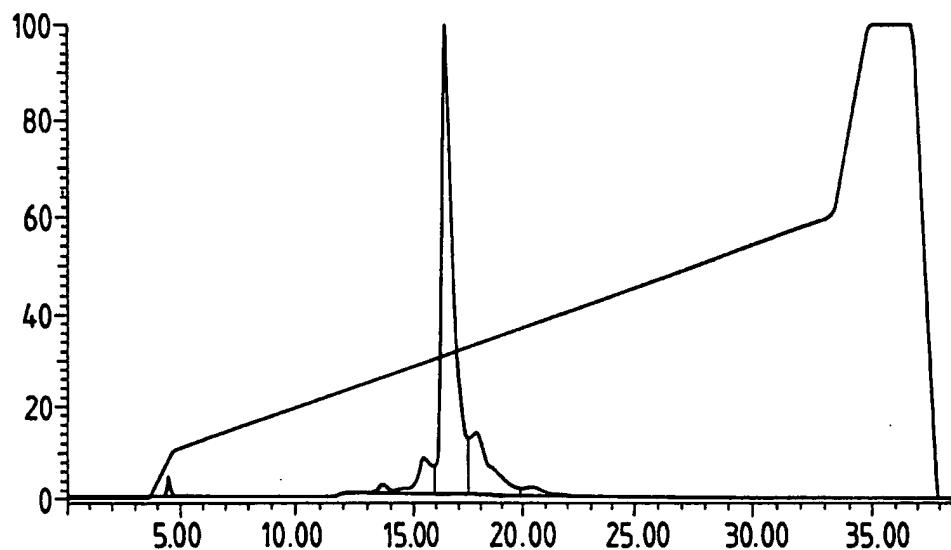
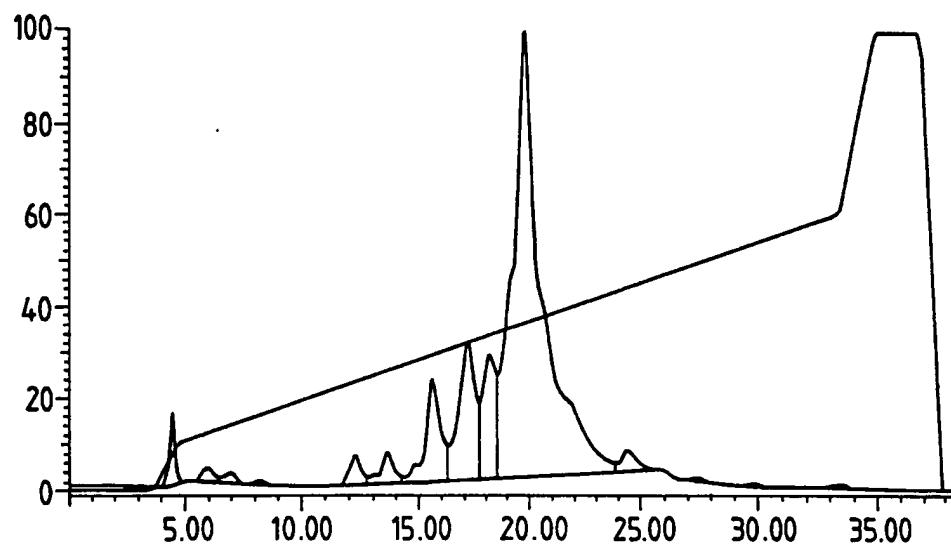


Fig.1b.



2/5

Fig.2a.

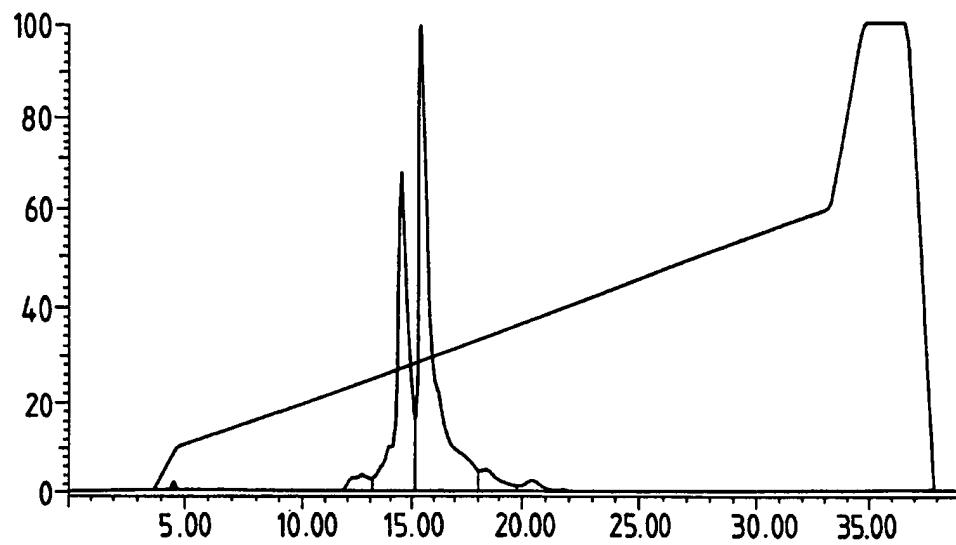
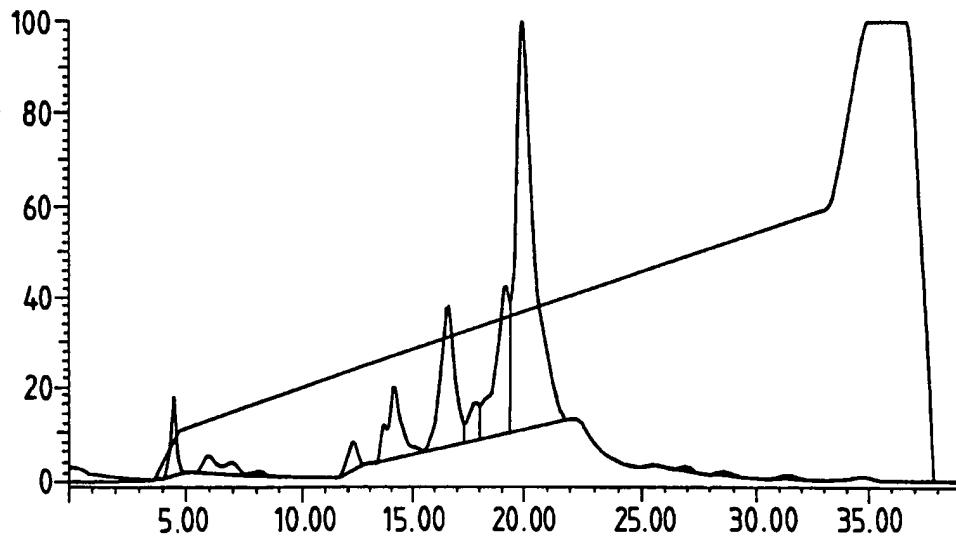


Fig.2b.



3/5

Fig.3a.

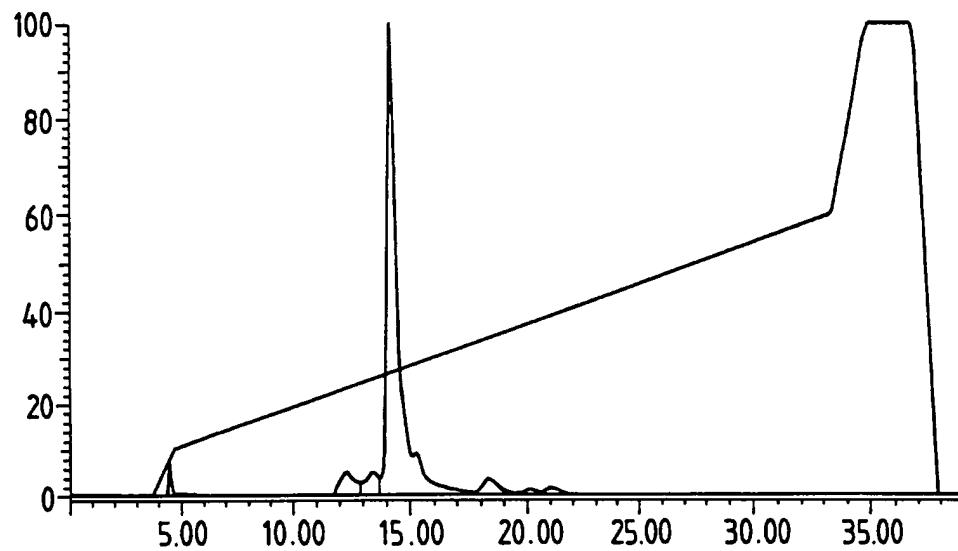
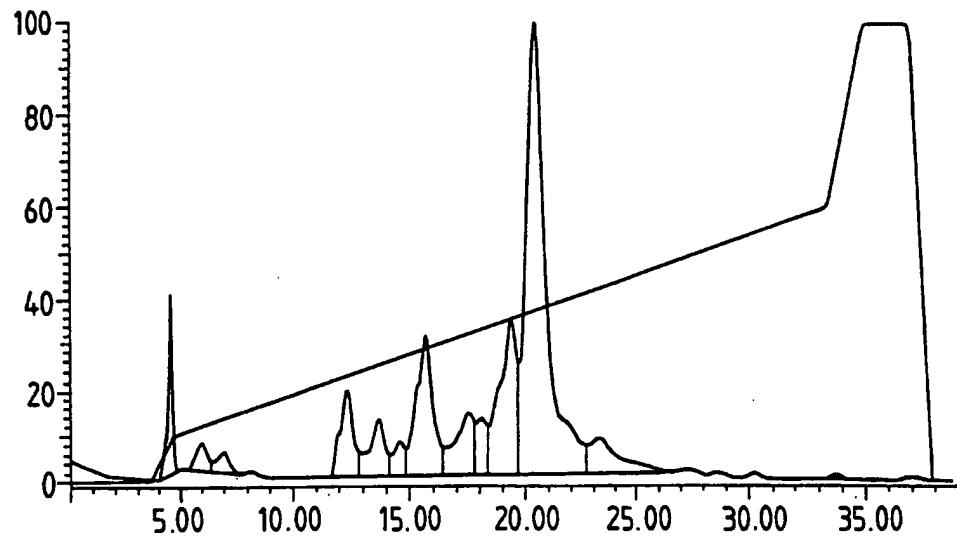


Fig.3b.



4/5

Fig.4a.

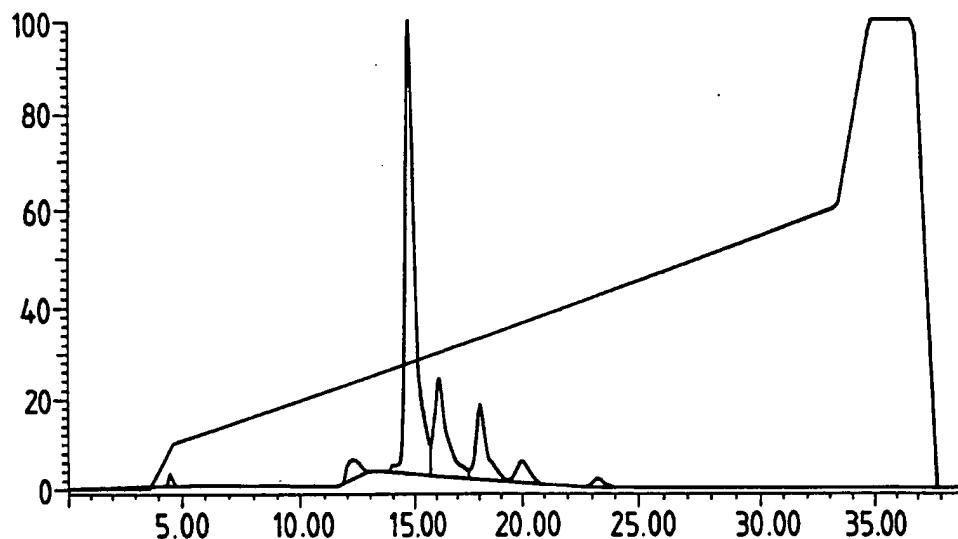
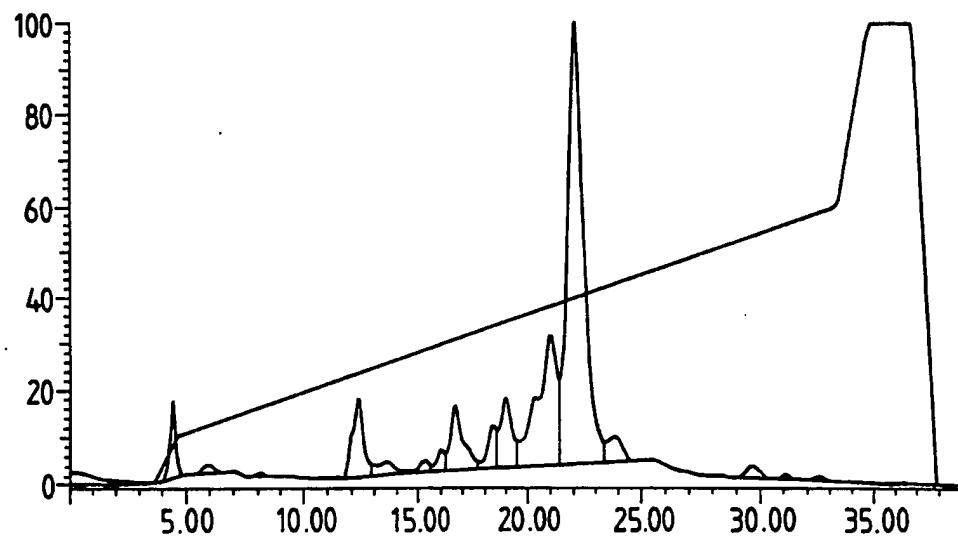


Fig.4b.



5/5

Fig.5a.

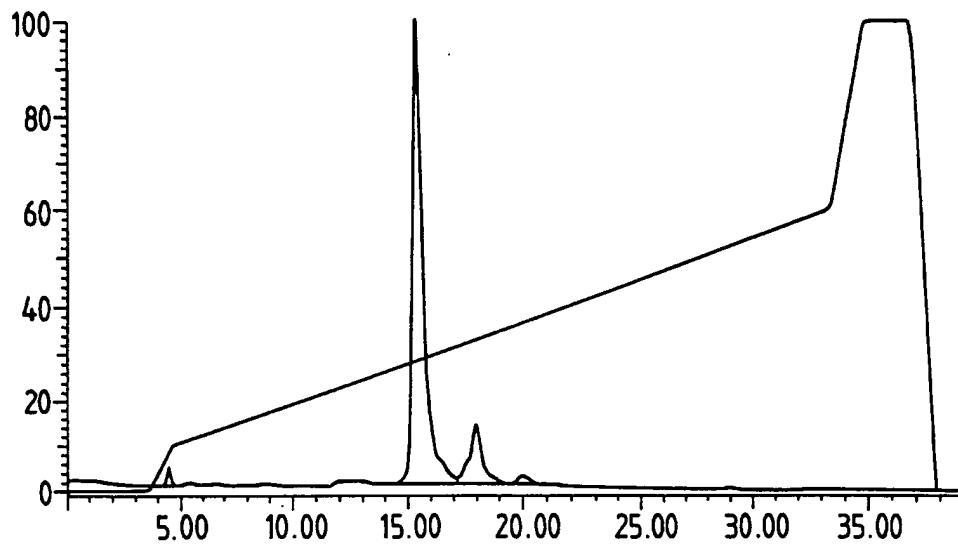
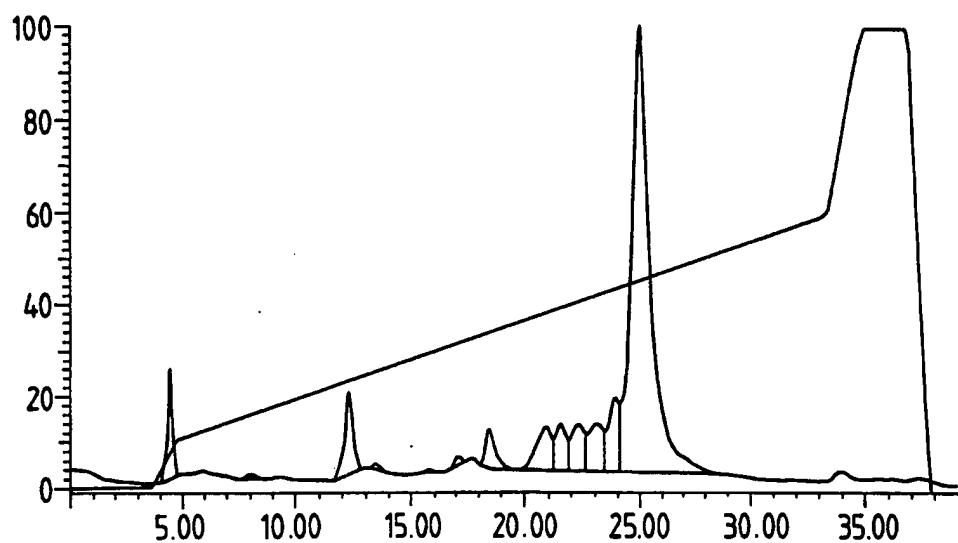


Fig.5b.



## INTERNATIONAL SEARCH REPORT

Int.	Application No
PCT/GB 94/02053	

A. CLASSIFICATION OF SUBJECT MATTER		
IPC 6	C07H21/00	C12Q1/68

According to International Patent Classification (IPC) or to both national classification and IPC
---

B. FIELDS SEARCHED
--------------------

Minimum documentation searched (classification system followed by classification symbols)
---

IPC 6 C07H C12Q
-----------------

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched
---

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)
--

C. DOCUMENTS CONSIDERED TO BE RELEVANT
--

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	EP,A,0 411 186 (ABBOTT LABORATORIES) 6 February 1991 see column 3, line 39 - line 53 see column 6, line 40 - line 46 ----	1-3
X	NUCLEIC ACIDS RESEARCH., vol.20, no.14, 25 July 1992, ARLINGTON, VIRGINIA US pages 3551 - 3554 A.SKERRA see the whole document ----	1-3,6,8
X	EP,A,0 546 761 (BECTON, DICKINSON & COMPANY) 16 June 1993 see page 7, line 9 - line 12 ----	1-3,6-9
		-/-

<input checked="" type="checkbox"/> Further documents are listed in the continuation of box C.
--

<input checked="" type="checkbox"/> Patent family members are listed in annex.
--

\* Special categories of cited documents :

- \*A\* document defining the general state of the art which is not considered to be of particular relevance
- \*B\* earlier document but published on or after the international filing date
- \*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- \*O\* document referring to an oral disclosure, use, exhibition or other means
- \*P\* document published prior to the international filing date but later than the priority date claimed

\*T\* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

\*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

\*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

\*&\* document member of the same patent family

1

Date of the actual completion of the international search
---

Date of mailing of the international search report
--

25 November 1994
------------------

21.12.94
----------

Name and mailing address of the ISA European Patent Office, P.B. 5818 Patentiaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo nl Fax: (+31-70) 340-3016
---

Authorized officer
--------------------

De Kok, A
-----------

## INTERNATIONAL SEARCH REPORT

Int.	Int'l Application No
PCT/GB 94/02053	

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	DE,A,41 10 085 (BOEHRINGER INGELHEIM INTERNATIONAL GMBH) 1 October 1992 see page 3, line 20 - line 30 see page 3, line 41 - line 43 ---	1-3,8,9
X	WO,A,92 20703 (O.BUCHARDT ET AL.) 26 November 1992 cited in the application * whole document, especially page 6, lines 23-27 and page 20, lines 35-36 *	1-5
A	EP,A,0 224 126 (THE UNIVERSITY OF CALGARY) 3 June 1987 see the whole document ---	10-13
A	TRENDS IN BIOTECHNOLOGY, vol.11, September 1993, CAMBRIDGE GB pages 384 - 386 O.BUCHARDT ET AL. see the whole document -----	1-13

## INTERNATIONAL SEARCH REPORT

Information on patent family members

Inten	al Application No
PCT/GB 94/02053	

Patent document cited in search report	Publication date	Patent family member(s)		Publication date
EP-A-0411186	06-02-91	NONE		
EP-A-0546761	16-06-93	AU-A- CA-A-	2844792 2083740	17-06-93 12-06-93
DE-A-4110085	01-10-92	NONE		
WO-A-9220703	26-11-92	AU-A- AU-A- CA-A- WO-A- EP-A- EP-A- JP-T- JP-T- NO-A- NO-A-	1880692 1884392 2109320 9220702 0586474 0586618 6509063 6506945 934122 934235	30-12-92 30-12-92 25-11-92 26-11-92 16-03-94 16-03-94 13-10-94 04-08-94 11-01-94 20-01-94
EP-A-0224126	03-06-87	JP-A-	63219380	13-09-88